





1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/856,070A

DATE: 03/17/2003 TIME: 15:41:36

Input Set: N:\vernette\US09856070A.raw
Output Set: N:\CRF4\03172003\I856070A.raw

```
1 (110 APPLICABT: Holms, Expert D.
      2 <120 - TITLE OF INVENTION: Regulatory/Unfolding Peptides of Ezrin
      3 (130) FIRE PEFERENCE: GUE-67
C--> 4 <140> CURRENT APPLICATION NUMBER: US/09/856,070A
C--> 5 <141> CURRENT FILING DATE: 2000-08-16
      6 (150) PETOR APPLICATION NUMBER: PCT/GB00/03566
      1 - (111) - PF : 05 FILING DAPE: 2000-09-15
      * 4150 PEIOR APPLICATION NUMBER: 9921881.0
                                                                  ENTERED
      9 (11) FF10E FILING DAME: 1999-09-17
     10 K160 KNUMBER OF SEQ ID NOS: 28
     11 (17) SCFTWARE: Patentin version 3.1
     15 MAD RESENTED NO: 1
     14 (211) BENGTH: 32
     15 KULLI TYPE: PRI
     16 -021 st OFCANISM: Artificial Sequence
     17 KOLOG PEATURE:
     1 - C.A. St. OTHER INFORMATION: Hepreseptor peptide
     1 - MIDU SEQUENCE: 1
             - Ala Arg Glu Glu Lys His Glr Lys Gln Leu Glu Arg Gln Gln Leu Glu
                                                    10
              Th: Gla Lye Lys Arg Arg Glu Thr Val Glu Arg Glu Lys Glu Gln Met
     1.5
                                                .25
     26 (210 EEQ 11 DO: 2
26 (211) LEUGTE: 34
     NV KNILL TYPE: PRT
     THE UNISH THE WANTEM: Artificial Sequence
     .. 4 Mill OF BEATTEE:
     30 th St OTHER INFORMATION: Hepreceptor peptide
     NO ON OF FEATURE:
     M. HOR OF NAME / KEY: MISC FEATURE
     ES \times Child Domation: (0.4)^{-}. (14)^{-}
     34 -12733 OTHER INFORMATION: Xaa = Tyr(P)
     FF GEOGRAFIENCE: 2
W - - > 36
              Met Arg Glu Lys Glu Glu Leu Met Leu Arg Leu Gln Asp Xaa Glu Glu
                                                    1.0
              Ly: Thr Lys Lys Ala Glu Arg Glu Leu Ser Glu Gln Ile Gln Arg Ala
     - }
                           20
     39
     4 (
             Le. Gln
     42 HISCO SEQ ID NO: 5
     43 KINDID LENGTH: 5
     44 KHIZO TYPE: PET
```

46 <210> FEATURE:

45 <2100 OEGANISM: Artificial Sequence

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Taput Set : N:\vernette\US09856070A.raw
Cutput Set: N:\CRF4\03172003\I856070A.raw

```
47 <2235 OTHER INFORMATION: Hepreceptor peptide
48 <40% · SEQUENCE: 3
        Th: Gua Lys Lys Arg
.1 +
5:)
        1
5.1 -1210 - SEP IT NO: 4
SK SOLL S LENGTH: 9
14 -0012 - TYPE: PHT
55 0013 0 0RGANISM: Artificial Sequence
14 HOLD FRATTE:
* / ALL R - OTHER INFORMATION: Hepredeptor peptide
No (400 - SEQUENCE: 4
4, 4
      Thr Glu Lys Lys Arg Arg Glu Thr Val
(1)
                          5
war ward to SEQ TH NO: 5
www.mille.hength: 11
64 -CHIC - TYPE: PRT
65 Miles ONSAMISM: Artificial Sequence
this con PRATIFE:
67 - WL: - OTHER INFORMATION: Hepredeptor peptide
6- - 400 - SEQUENCE: 5
18 Thr Glu Lys Lys Arg Arg Glu Thr Val Glu Arg
        1
                         1,
7. SUDDA SEQ II 100: 6
73 -0.1110 LENGTH: 5
74 < 1.1.  TYPE: PM
T(-1,1) = 0 \oplus AMSM: Artificial Sequence
76 KIND FEATURE:
// WILE OTHER INFORMATION: Hepreceptor peptide
79 - 49 DESEMBLICE: 6
7.4
        Lys Lys Arg Arg Glu
20
        1
M. H. 100 SEQ 10 DO: 7
84 HOLLIN LENGTH: 8
*4 *1.1LP TYPE: PFT
85 CLIBS CHUANISM: Artificial Sequence
HO STATULE:
FF RULES OTHER INFOFMATION: Hepreceptor peptide
For Education SPOURNCE: 7
(p. 3)
   Lys Lys Arg Arg Glu Thr Val Glu
40
                          5
        1
#. 40100 SEQ H NO: 8
93 -0.000 DENGTH: 10
94 -0 DI TYPE: PHT
9' - C.13: OFGANISM: Artificial Sequence
'me sull'us FEATURE:
9\% < \!\! 20.30 \cdot 00 \mbox{MEF} INFCFMATION: Hepreceptor peptide
95 <4005 SEQUENCE: 8
9+ Lys Lys Arg Arg Glu Thr Val Glu Arg Glu
100
                      5
        ]
```

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PATENT APPLICATION: US/09/856,070A

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Input Set: N:\vernette\US09856070A.raw Output Set: N:\CRF4\03172003\I856070A.raw

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100 <210> SEO ID NO: 9
1-1 - (211> LEMGTH: 11
194 -021. N TYPE: EPT
105 - 1213 - ORSANISM: Artificial Sequence
100 HRRD - FEATURE:
187 - Char OTHER INFORMATION: Hepreceptor peptide
10% -040% - SEQUENCE: 9
154
          Lys Lys Ard Ard Glu Fnr Val Glu Ard Glu Lys
110
                           1,
          1
112 - Mio - SEQ ID NO: 10
11: 0.011 - LENGTH: 12
114 HOLD FIRE: EMT
11^{\rm t} - 0.01 \, {\rm t} + 0 {\rm H/MADDSM:} Artifitial Sequence
110 Hillion PEATURE:
117 HARBY OTHER INFOSMATION: Hepreceptor peptide
115 -0400 - SEQUENCE: 10
119 Lys Lys Ard Ard Glu Thr Val Glu Ard Glu Lys Glu
         1
1. .
                          ٠,
100 BINDS SEQ ID NO: 11
100 HOLD - BENGTH: 5
1.4 dull - TYPE: FET
1.5 RMIN ONGAMISM: Artificial Sequence
1.6 Same FRATURE:
12 1 - Children OTHER INFORMATION: Hepredeptor peptide
118 - HOW - SEQUENCE: 11
          Lys Ang Ang Glu Thr Val Glu Ang
1. 🔞
120
          1
150 -0.000- SEQ OF NO: 12
13: HELLE DENGTH: 16
134 Hills TYPE: FAT
145 - Color OFFANISM: Antificial Sequence
1 % SULTON FEATURE:
1:7 H. .. S. OTHER INFORMATION: Hepreceptor peptide
138 -04000 SEQUENCE: 12
    Dys Ang Ang Glu Thr Val Glu Ang Glu Dys
1 5 10
1 . . .
140 -0100 SEQ ID NO: 13
143 -00110 LENGTH: 11
144 H. D.D. DYPE: FHI
145 -LIBE OF GANUSM: Artificial Sequence
146 -LICE FEATURE:
147 - MINGS OTHER INFORMATION: Hepreceptor peptide
148 - 4000 SEQUENCE: 13
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1.1.+
150
                                                 10
150 -0.100 SEQ ID NO: 14
155 -0.110 LFNGTH: 5
154 < 170 TYPE: PET
155 < 15> OFGANISM: Artificial Sequence
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## RAW SEQUENCE LISTING

FATENT APPLICATION: US/09/856,070A TIME: 15:41:36

DATE: 03/17/2003

Imput Set : N:\vernette\US09856070A.raw
Cutput Set: N:\CRF4\03172003\I856070A.raw

```
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1 7 K. . OTHER INFORMATION: Hepreceptor peptide
1'8 <4 -- SEQUENCE: 14
1: 3
          Arr Arg Glu Thr Val
1 . . .
160 0010 SEQ 10 NO: 15
1. . . II - LEUGTH: 9
1.4 -Clastype: PMT
163 HOURS ARTIFICIAL Sequence
146 H. J. PEARTHE:
1.7 dans OTHER INFORMATION: Hepreceptor peptide
16% (GEO - SEQUENCE: 15
    Argellu Thr Val Glu Argellu Lys Glu
1 \cdot +
17:
14. 4.45 SEQ II 10: 16
175 40 HI - DENGTH: 5
174 - 117 - 19PE: PET
12 GD1 - OBGANISM: Artificial Sequence
176 HATCH PEATURE:
1/1 RE. 4 OTHER INFORMATION: Hepreceptor peptide
178 KAUCH SEQUENCE: 16
1 ***
          Gla Ara Glu Lys Glu
180
          1
1%: 0010 - SEQ ID NO: 17
1% - 0.11 | DENGTH: 14
1%4 | 0.11 | TYPE: FET
1%4 | 0.11 | OF-ANISM: Artificial Sequence
186 - C. Just FRATURE:
1c7 to Add OTHER INFORMATION: Hepreceptor peptide
1-8 (400 - SEQUENCE: 17
1-4 31: Arg Glu Lys Glu Gln Met Met Arg Glu Lys Glu Glu Leu
1
191 -02175 SEQ II NO: 18
194 -COLI - LEDGTH: 8
194 BUILDS TYPE: ERI
135 C. 13 · OF CANDEM: Artificial Sequence
100 C. NOW PRATURE:
140 A.A.B. OTHER INFORMATION: Hepreceptor peptide
1 % (4000 SECUENCE: 18
199 - Byr Glu Glu Leu Met
2000
         1
201 0 100 SEC 10 NO: 19
200 - 1.110 LFNGTH: 13
204 HOLLS TYPE: FRT
201 - C. LES OEGANISM: Artificial Sequence
200 -01100- PHATURE:
207 KIRE INFOFMATION: Hepreceptor peptide
208 <400> SEQUENCE: 19
209 Ly: Glu Glu Leu Met Leu Arg Leu Gln Asp Tyr Glu Glu
```

DATE: 03/17/2003

TIME: 15:41:36

## Imput Set : N:\vernette\US09856070A.raw Output Set: N:\CRF4\03172003\1856070A.raw 210 1 1.0 21.1 <210> SEC ID NO: 20 .11 < .11> DENGTH: 13 .14 <13> TYPE: PRI .15 G.178 OR WANTEM: Artificial Sequence .16 CLR: FEATURE: 717 HOURS OTHER INFORMATION: Hepreceptor peptide MIN WHITE FEATURE: 714 4.01 - NAME KEY: MISC FEATURE $220 - 1.2. = 10 \text{ ATION: } (11) \dots (11)$ . . . . . . . . OTHER INFORMATION: Xaa = Tyr(P) ... Gus - AROUNCE: ..O Lys Glu Glu Leu Met Leu Arg Leu Gln Asp Xaa Glu Glu W--> 223 ...... ι, DIV KNICH CER IN 199: 31 ZZ 1 KOZZ K BENGTH: 12 JUN KLID KITTER PRI ... Artificial Sequence . SHOULD FEARIFE: AND AN AND HER INFORMATION: Hepreceptor peptide I. 440 - PEQUENCE: II - Gla Gla Les Met Lea Arg Lea Gln Asp Tyr Gla Gla . 1 1.0 Ė 5.20 (21) (780 H NO: 32 LE7 - M.11 - LENGTH: 12 Describe Type: Phr 13 · 4.14 OF PANISM: Artificial Sequence 140 HAROS FFATURE: 141 - 1187 - OTHER INFORMATION: Hepreceptor peptide 147 HIRL FRATURE: 14 % SUBTO MAME/FEY: MISC FEATURE $244 \times 32.3 \times LOMATION: (10) \overline{.}. (10)$ .Mf +0.150+ OTHER INFORMATION: Xaa = Tyr(P) 146 HADON SEQUENCE: 11 W--> 247Glu Glu Leu Met Leu Arg Leu Gln Asp Xaa Glu Glu 2.18 750 -0710 - 8E0 IF 10: 13 151 -1110 LENGTH: 11 LAN HUIND TYPE: PHT 1855 ALIBE (FRANISM: Artificial Sequence .54 <2000 FEATUFE: MAN WARR OTHER INFORMATION: Hepreceptor peptide JER HADDE SEGUENCE: 25 1:7 Glu Leu Met Leu Arg Leu Gln Asp Tyr Glu Glu . 18 10 .00 00 100 SEQ IL NO: 04 161 <0.110 SFUGTH: 11 1.621 KHILD TYPE: PRT 263 < 15> OFCANISM: Artificial Sequence

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/856,070A

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 03/17/2003 PATENT APPLICATION: US/09/856,070A

TIME: 15:41:37

Input Set : N:\vernette\US09856070A.raw Output Set: N:\CRF4\03172003\1856070A.raw

## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:0; Xaa Pos. 14 Seq#:110; Maa Pos. 11 Seq#:22; Maa Pos. 10 Seq#:24; Maa Pos. 9 Seq#:27; Maa Pos. 3

# VERIFICATION SUMMARY

DATE: 03/17/2003 PATENT APPLICATION: US/09/856,070A TIME: 15:41:37

input Set : N:\vernette\US09856070A.raw Output Set: N:\CRF4\03172003\1856070A.raw

L:4 M:270 C: Current Application Number differs, Wrong Format L:5 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:36 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0 L:223 M:341 W: (46) "n" or "Maa" used, for SEQ ID#: 0 after pos.:0 L:347 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0 L:371 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0 L:305 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:0